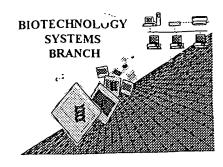
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/765,231
Source:	OIPE
Date Processed by STIC:	7/5/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old-rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING

DATE: 07/05/2001

PATENT APPLICATION: US/09/765,231

TIME: 11:35:21

Input Set : N:\paola\09765231.txt

Output Set: N:\CRF3\07032001\I765231.raw

Does Not Comply Corrected Diskette Needed

225

2 <110> APPLICANT: Searle/Monsanto 3

Phippard, Deborah

Vasanthakamur, Geetha

Dotson, Stanton

Ma, Xiao-Jun

<120> TITLE OF INVENTION: Osteoarthritis tissue-derived nucleic acids, polypeptides,

vectors, and cells

11 <130> FILE REFERENCE: SO-3221 PR

13 <140> CURRENT APPLICATION NUMBER: US/09/765,231

13 <141> CURRENT FILING DATE: 2001-01-18

13 <160> NUMBER OF SEQ ID NOS: 82

ERRORED SEQUENCES

3483 <210> SEQ ID NO: 82

3484 <211> LENGTH: 225

3485 <212> TYPE: DNA

3486 <213> ORGANISM: Homo sapiens

3488 <400> SEQUENCE: 82

3490 ggagaatgtg acatagattt gctggcacat gggtttccta tgagcaaacc ccagaattgg 60

3492 acacacgtat ctggtgctgc attggaatca tccgaaaaaa ccaaggcttg cattgcatat

180 3494 ctatctgctg tetgetgaag gageeetgte tgtgtgeeea aggaagtgae atcettgeea

for more ever

3496 agggctgtcc ctgttgcagg agatqaagga gccctgtcta tgtgc

E--> 3503/1

delite at md of file

7/5/01

<pre>< <210> <211> <212></pre>	6 2263 DNA	Resid
<213>	Homo sapiens	7/2227
<223> <400>	unsure at all n locations	
		L2227-1

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

MI

Use of n and/or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

C2217 / Lese, [2207 is "header only it las no

Besides (223)

VERIFICATION SUMMARY PATENT APPLICATION: US/09/765,231

DATE: 07/05/2001 TIME: 11:35:22

Input Set : N:\paola\09765231.txt

Output Set: N:\CRF3\07032001\I765231.raw

```
L:13 M:270 C: Current Application Number differs, Replaced Current Application No
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:251 M:283 W: Missing Blank Line separator, <400> field identifier
L:255 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:6
L:255 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:6
L:255 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:6
L:255 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:426 M:283 W: Missing Blank Line separator, <400> field identifier
L:430 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:9
L:430 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
L:430 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
L:430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:432 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:9
L:432 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:9
L:432 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:9
L:432 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:708 M:283 W: Missing Blank Line separator, <400> field identifier
L:756 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:756 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:14
L:756 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:14
L:756 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:1139 M:283 W: Missing Blank Line separator, <400> field identifier
L:1141 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:21
L:1141 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:21
L:1141 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:21
L:1141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:1432 M:283 W: Missing Blank Line separator, <400> field identifier
L:1458 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:27
L:1458 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:27
L:1458 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:27
L:1458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1635 M:283 W: Missing Blank Line separator, <400> field identifier
L:1641 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:30
L:1641 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:30
L:1641\ M:258\ W: Mandatory Feature missing, <222> not found for SEQ ID#:30
L:1641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:2386 M:283 W: Missing Blank Line separator, <400> field identifier
L:2424 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:41
L:2424 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:41
L:2424 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:41
L:2424 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:2493 M:283 W: Missing Blank Line separator, <400> field identifier
L:2495 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:42
L:2495\ M:258\ W: Mandatory Feature missing, <221> not found for SEQ ID#:42
L:2495 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:42
L:2495 \text{ M}:341 \text{ W}: (46) \text{ "n" or "Xaa" used, for SEQ ID$#:42}
L:3080 M:283 W: Missing Blank Line separator, <400> field identifier
L:3082 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:61
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/765,231

Input Set : N:\paola\09765231.txt

Output Set: N:\CRF3\07032001\1765231.raw

DATE: 07/05/2001

TIME: 11:35:22

L:3082 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:61 L:3082 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:61 L:3082 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 L:3084 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:61 L:3084 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:61 L:3084 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:61 L:3084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 L:3411 M:283 W: Missing Blank Line separator, <400> field identifier L:3415 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:79 L:3415 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:79 L:3415 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:79 L:3415 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 L:3417 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:79 L:3417 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:79 L:3417 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:79 $L\!:\!3417$ M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 L:3435 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:79 L:3435 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:79 L:3435 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:79 L:3435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 L:3446 M:283 W: Missing Blank Line separator, <400> field identifier L:3448 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:80 L:3448 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:80 L:3448 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:80 L:3448 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 L:3450 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:80 L:3450 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:80 L:3450 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:80 L:3450 M:341 W: (46) "n" or "Xaa" used, for SEQ ID\$\$#:80\$L:3503 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:225 SEQ:82